SEQUENCE LISTING

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Kumagai, Hidetoshi

<120> MAST CELL-DERIVED MEMBRANE PROTEINS

<130> 14875-142US1

<150> PCT/JP2003/013921
<151> 2003-10-30

<150> JP 2002-316680
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<150> JP 2002-354165
<151> 2002-12-05

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<170> PatentIn version 3.1

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cctgccggtg acccgtgtgt gggagaa atg acc caa ctg gcc tca gct gtg tgg 174
Met Thr Gln Leu Ala Ser Ala Val Trp
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ctg ccc acg ctg ttg ctg ctg ctg ctg ctt ttt tgg ctt cca ggc tgt 222
Leu Pro Thr Leu Leu Leu Leu Leu Leu Phe Trp Leu Pro Gly Cys
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gtc cct ctg cat ggt ccc agc acc atg aca gga agt gtg ggt caa tcc 270
Val Pro Leu His Gly Pro Ser Thr Met Thr Gly Ser Val Gly Gln Ser
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ctg agt gtg tcg tgt cag tat gag gag aaa ttt aag act aag gac aaa 318
Leu Ser Val Ser Cys Gln Tyr Glu Glu Lys Phe Lys Thr Lys Asp Lys
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tac tgg tgc aga ggg tca ctt aag gta ctg tgc aaa gat att gtc aag 366
Tyr Trp Cys Arg Gly Ser Leu Lys Val Leu Cys Lys Asp Ile Val Lys
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acc agc agc tca gaa gaa gct agg agt ggc aga gtg acc atc agg gac 414

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Thr Ser Ser Ser Glu Glu Ala Arg Ser Gly Arg Val Thr Ile Arg Asp
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cat cca gac aac ctc acc ttc aca gtg acc tat gag agc ctc acc ctg      462
His Pro Asp Asn Leu Thr Phe Thr Val Thr Tyr Glu Ser Leu Thr Leu
90                  95                  100                 105

gat gat gca gac acc tac atg tgt gcg gtg gat ata cca ttt ttc aat      510
Asp Asp Ala Asp Thr Tyr Met Cys Ala Val Asp Ile Pro Phe Phe Asn
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gcc ccc ttg ggg ctc gat aag tac ttc aag att gaa ttg tct gtg gtt      558
Ala Pro Leu Gly Leu Asp Lys Tyr Phe Lys Ile Glu Leu Ser Val Val
            125                 130                 135

cca agt gag gac cca gtt tca tct cca gga cca aca cta gag aca cct      606
Pro Ser Glu Asp Pro Val Ser Ser Pro Gly Pro Thr Leu Glu Thr Pro
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gtg gtg tcc acc agt ctg cct acc aag ggt ccc gcc cta gga tcc aac      654
Val Val Ser Thr Ser Leu Pro Thr Lys Gly Pro Ala Leu Gly Ser Asn
    155                 160                 165

aca gag gac cgc cgt gag cat gac tat tcc cag ggc ttg agg ctc cca      702
Thr Glu Asp Arg Arg Glu His Asp Tyr Ser Gln Gly Leu Arg Leu Pro
170                 175                 180                 185

gcg ctg ttg tct gtg tta gct ctc ctg ctg ttt ctg ttg gtg ggg aca      750
Ala Leu Leu Ser Val Leu Ala Leu Leu Leu Phe Leu Leu Val Gly Thr
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tct ctg ctg gcc tgg agg atg ttc cag aag cgg ctg gtc aaa gct gat      798
Ser Leu Leu Ala Trp Arg Met Phe Gln Lys Arg Leu Val Lys Ala Asp
            205                 210                 215

agg cat cca gag ctg tcc cag aac ctc aga cag gct tct gag cag aat      846
Arg His Pro Glu Leu Ser Gln Asn Leu Arg Gln Ala Ser Glu Gln Asn
        220                 225                 230

gag tgc cag tat gtg aat ttg cag ctg cac acg tgg tct ctg agg gaa      894
Glu Cys Gln Tyr Val Asn Leu Gln Leu His Thr Trp Ser Leu Arg Glu
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gag ccg gtg cta cca agt cag gta gaa gtg gtg gaa tat agc aca ttg      942
Glu Pro Val Leu Pro Ser Gln Val Glu Val Val Glu Tyr Ser Thr Leu
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gca tta ccc cag gaa gag ctt cac tat tca tcc gtg gca ttc aac tcc      990
Ala Leu Pro Gln Glu Glu Leu His Tyr Ser Ser Val Ala Phe Asn Ser
                270                 275                 280

cag agg cag gat tct cac gcc aat gga gat tct ctt cat caa cct cag     1038
Gln Arg Gln Asp Ser His Ala Asn Gly Asp Ser Leu His Gln Pro Gln
            285                 290                 295

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Thr Met Thr Gly Ser Val Gly Gln Ser Leu Ser Val Ser Cys Gln Tyr
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Glu Glu Lys Phe Lys Thr Lys Asp Lys Tyr Trp Cys Arg Gly Ser Leu
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Lys Val Leu Cys Lys Asp Ile Val Lys Thr Ser Ser Ser Glu Glu Ala
65                  70                  75                  80

Arg Ser Gly Arg Val Thr Ile Arg Asp His Pro Asp Asn Leu Thr Phe
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Thr Val Thr Tyr Glu Ser Leu Thr Leu Asp Asp Ala Asp Thr Tyr Met
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Cys Ala Val Asp Ile Pro Phe Phe Asn Ala Pro Leu Gly Leu Asp Lys
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Tyr Phe Lys Ile Glu Leu Ser Val Val Pro Ser Glu Asp Pro Val Ser
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Ser Pro Gly Pro Thr Leu Glu Thr Pro Val Val Ser Thr Ser Leu Pro
145                 150                 155                 160

Thr Lys Gly Pro Ala Leu Gly Ser Asn Thr Glu Asp Arg Arg Glu His
                165                 170                 175

Asp Tyr Ser Gln Gly Leu Arg Leu Pro Ala Leu Leu Ser Val Leu Ala
            180                 185                 190

Leu Leu Leu Phe Leu Leu Val Gly Thr Ser Leu Leu Ala Trp Arg Met
        195                 200                 205

Phe Gln Lys Arg Leu Val Lys Ala Asp Arg His Pro Glu Leu Ser Gln
    210                 215                 220

Asn Leu Arg Gln Ala Ser Glu Gln Asn Glu Cys Gln Tyr Val Asn Leu
225                 230                 235                 240

Gln Leu His Thr Trp Ser Leu Arg Glu Glu Pro Val Leu Pro Ser Gln
                245                 250                 255

Val Glu Val Val Glu Tyr Ser Thr Leu Ala Leu Pro Gln Glu Glu Leu
            260                 265                 270

His Tyr Ser Ser Val Ala Phe Asn Ser Gln Arg Gln Asp Ser His Ala
        275                 280                 285

Asn Gly Asp Ser Leu His Gln Pro Gln Asp Gln Lys Ala Glu Tyr Ser
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Glu Ile Gln Lys Pro Arg Lys Gly Leu Ser Asp Leu Tyr Leu
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tct cag gtc cca ggc tgt gtc cca ctg cat ggc ccc agc act atc aca       96
Ser Gln Val Pro Gly Cys Val Pro Leu His Gly Pro Ser Thr Ile Thr
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ttc aag act aag gac aaa ttc tgg tgc aga ggg tca ctg aag gta ctc      192
Phe Lys Thr Lys Asp Lys Phe Trp Cys Arg Gly Ser Leu Lys Val Leu
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tgt aaa gat att gtc aag acc agc agc tca gaa gaa gtt agg aat ggc      240
Cys Lys Asp Ile Val Lys Thr Ser Ser Ser Glu Glu Val Arg Asn Gly
65                  70                  75                  80

cga gtg acc atc agg gac cat cca gac aac ctc acc ttc aca gtg acc      288
Arg Val Thr Ile Arg Asp His Pro Asp Asn Leu Thr Phe Thr Val Thr
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tat gag agc ctc acc ctg gag gat gca gac acc tac atg tgt gcg gtg      336
Tyr Glu Ser Leu Thr Leu Glu Asp Ala Asp Thr Tyr Met Cys Ala Val
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gat ata tca ctt ttt gat ggc tcc ttg ggg ttc gat aag tac ttc aag      384
Asp Ile Ser Leu Phe Asp Gly Ser Leu Gly Phe Asp Lys Tyr Phe Lys
        115                 120                 125

att gag ttg tct gtg gtt cca agt gag gac cca gtc aca ggt tcg agc      432
Ile Glu Leu Ser Val Val Pro Ser Glu Asp Pro Val Thr Gly Ser Ser
    130                 135                 140

ctt gag agt ggt aga gat atc ctg gaa tcc ccc aca tcc tca gtt ggg      480
Leu Glu Ser Gly Arg Asp Ile Leu Glu Ser Pro Thr Ser Ser Val Gly
145                 150                 155                 160

cac act cat ccc agt gtg acc aca gat gac aca att cct gct ccc tgc      528
His Thr His Pro Ser Val Thr Thr Asp Asp Thr Ile Pro Ala Pro Cys
                165                 170                 175

cct cag cct cgg tct ctt cgg agc agc ctc tac ttc tgg gtc ctg gtg      576
Pro Gln Pro Arg Ser Leu Arg Ser Ser Leu Tyr Phe Trp Val Leu Val
            180                 185                 190

tct ctg aag ttg ttc ctg ttc ctg agc atg ctt ggt gct gtc ctc tgg      624
Ser Leu Lys Leu Phe Leu Phe Leu Ser Met Leu Gly Ala Val Leu Trp
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gtg aac agg cct cag agg tgc tct ggg gga agc agc act cag ccc tgt      672
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Gly Ala Val Gly Glu Ser Leu Ser Val Ser Cys Gln Tyr Glu Glu Lys
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Phe Lys Thr Lys Asp Lys Phe Trp Cys Arg Gly Ser Leu Lys Val Leu
    50                  55                  60

Cys Lys Asp Ile Val Lys Thr Ser Ser Ser Glu Glu Val Arg Asn Gly
65                  70                  75                  80

Arg Val Thr Ile Arg Asp His Pro Asp Asn Leu Thr Phe Thr Val Thr
                85                  90                  95

Tyr Glu Ser Leu Thr Leu Glu Asp Ala Asp Thr Tyr Met Cys Ala Val
            100                 105                 110

Asp Ile Ser Leu Phe Asp Gly Ser Leu Gly Phe Asp Lys Tyr Phe Lys
        115                 120                 125

Ile Glu Leu Ser Val Val Pro Ser Glu Asp Pro Val Thr Gly Ser Ser
    130                 135                 140

Leu Glu Ser Gly Arg Asp Ile Leu Glu Ser Pro Thr Ser Ser Val Gly
145                 150                 155                 160

His Thr His Pro Ser Val Thr Thr Asp Asp Thr Ile Pro Ala Pro Cys
                165                 170                 175

Pro Gln Pro Arg Ser Leu Arg Ser Ser Leu Tyr Phe Trp Val Leu Val
            180                 185                 190

Ser Leu Lys Leu Phe Leu Phe Leu Ser Met Leu Gly Ala Val Leu Trp
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Tyr Glu Asn Gln
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